

REMARKS

A Transmittal of Proposed Drawing Corrections and New Formal Drawings is being filed concurrently. As originally filed, certain amino acid residues in Figure 2 were framed in black. In the Proposed Drawing Correction submitted concurrently herewith, these residues are double-underlined instead. The specification is being amended accordingly to be consistent with the corrected figure.

Applicants' Attorney respectfully requests consideration and entry of the present amendment.

Respectfully submitted,

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Complete nucleotide sequence of IP10/MigR (MLRA) cDNA

CCAACCACAA GCACCAAAGC AGAGGGGCAG GCAGCACACC ACCCAGCAGC	-50- 60
CAGAGCACCA//GCCCAGCCAT GGTCTTTGAG GTGAGTGACC ACCAAGTGCT	-100- 120
AAATGACGCC GAGGTTGCCG//CCCTCCTGGA GAACTTCAGC TCTTCCTATG	-150- 180
ACTATGGAGA AAACGAGAGT GACTCGTGCT//GTACCTCCCC GCCCTGCCCA	-200- 240
CAGGACTTCA GCCTGAACTT CGACCGGGGCC TTCCTGCCAG//CCCTCTACAG	-250- 300
CCTCCTCTTT CTGCTGGGGC TGCTGGGCAA CGGCGCGGTG GCAGCCGTGC//	-300- 360
TGCTGAGCCG GCGGACAGCC CTGAGCAGCA CCGACACCTT CCTGCTCCAC	-350- 420
CTAGCTGTAG//CAGACACGCT GCTGGTGCTG ACACTGCCGC TCTGGGCAGT	-400- 480
GGACGCTGCC GTCCAGTGGG//TCTTTGGCTC TGGCCTCTGC AAAGTGGCAG	-450- 540
GTGCCCTCTT CAACATCAAC TTCTACGCAG//GAGCCCTCCT GCTGGCCTGC	-500- 600
ATCAGCTTTG ACCGCTACCT GAACATAGTT CATGCCACCC//AGCTCTACCG	-550- 660
CCGGGGGGCC CCGGGCCGCG TGACCCTCAC CTGCCTGGCT GTCTGGGGGC//	-600- 720
TCTGCCTGCT TTTGCCCTC CCAGACTTCA TCTTCCTGTC GGCCACCAC	-650- 780
GACGAGCGCC//TCAACGCCAC CCACTGCCAA TACAATTCC CACAGGTGGG	-700- 840
CCGCACGGCT CTGCGGGTGC//TGCAGCTGGT GGCTGGCTTT CTGCTGCCCC	-750- 900
TGCTGGTCAT GGCCTACTGC TATGCCCACA//TCCTGGCCGT GCTGCTGGTT	-800- 960
TCCAGGGGCC AGCGGCGCCT GCGGGCCATG CGGCTGGTGG//TGGTGGTCGT	-850- 1020
GGTGGCCTTT GCCCTCTGCT GGACCCCTA TCACCTGGTG GTGCTGGTGG//	-900- 1080
ACATCCTCAT GGACCTGGGC GCTTTGGCCC GCAACTGTGG CCGAGAAAGC	-950- 1140
AGGGTAGACG//TGGCCAAGTC GGTCACCTCA GGCCTGGGCT ACATGCACTG	-1000- 1200
CTGCCTCAAC CCGCTGCTCT//ATGCCTTTGT AGGGGTCAAG TTCCGGGAGC	-1050- 1260
GGATGTGGAT GCTGCTCTTG CGCCTGGGCT//GCCCCAACCA GAGAGGGCTC	-1100- 1320
CAGAGGCAGC CATCGTCTTC CCGCCGGGAT TCATCCTGGT//CTGAGACCTC	-1150- 1380
AGAGGCCTCC TACTCGGGCT TGTGAGGCCG GAATCCGGGC TCCCCTTTG//	-1200- 1440
CCCACAGTCT GACTTCCCCG CATTCCAGGC TCCTCCCTCC CTCTGCCGGC	-1250- 1500
TCTGGCTCTC//CCCAATATCC TCGCTCCCGG GACTCACTGG CAGCCCCAGC	-1300- 1560
ACCACCAGGT CTCCCGGGA//GCCACCCTCC CAGCTCTGAG GACTGCACCA	-1350- 1620
TTGCTGCTCC TTAGCTGCCA AGCCCCATCC//TGCCGCCCGA GGTGGCTGCC	-1400-
TGGAGCCCCA CTGCCCTTCT CATTTGGAAA CTAAAACTTC//ATCTTCCCCA	-1450-
AGTGCGGGGA GTACAAGGCA TGGCGTAGAG GGTGCTGCCC CATGAAGCCA//	-1500- 8
CAGCCCAGGC CTCCAGCTCA GCAGTGA CTG TGGCCATGGT CCCAAGACC	-1550-
TCTATATTTG//CTCTTTTATT TTTATGTCTA AAATCCTGCT TAAAACTTTT	-1600-
CAATAAACAA GATCGTCAGG//ACCTTTTTTT TTTTTTTTTT TTTTTTTTTT	-1650-
TTTTTTTTTT TTTTTTTTTT	1670

FIGURE 1

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 MVLEVDHQVLNDAEVAALLENFSSVYGENESDSCCTSPCEQDFSLNEIDRAEAEAVISLDEHGHGTCGGAAVAATSRRTALESSTDTFFKTHENVD
 TM 1 / TM 2
 90
 --99--

[illegible]

TM 5 / TM 6

Y
NATGQYN~~E~~EQVC-----ATAEVLQLVAGELEPPLVAAYCAIAHTEAVTVVSRGQRRLRAVRVVWVAFAICGFPRFVAVVDITMDLGNZARIGCG

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 TM 7
 RESRV~~VA~~KS~~VI~~SG~~CT~~M~~HCC~~NE~~LI~~MA~~VC~~K~~ER~~MM~~WT~~IR---L~~CC~~P~~NR~~Q~~CT~~Q~~RP~~SS~~SR~~RD~~SS~~W~~ET~~SEASYSGL
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FIGURE 2